

Serial Number: 09/234,290A

Changed a file from non-ASCII to ASCII

"Changed the margins in cases where the sequence text was 'wrapped' down to the next line

Edited a format error in the Current Application Data section, specifically:

Edited the Current Application Data section with the actual current number. The number input applicant was the prior application data; or other _____

Added the mandatory heading and subheadings for "Current Application Data".

Edited the "Number of Sequences" field. The applicant spelled out a number instead of using

Changed the spelling of a mandatory field (the headings or subheadings), specifically:

Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited

Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:

Corrected subheading placement. All responses must be on the same line as each subheading applicant placed a response below the subheading, this was moved to its appropriate place.

Inserted colons after headings/subheadings. Headings edited included: **ENTERED**

Deleted extra, invalid, headings used by an applicant, specifically:

Deleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename a page numbers throughout text; other invalid text, such as **RECEIVED**

Inserted mandatory headings, specifically:

Corrected an obvious error in the response, specifically: **NOV 14 2001**

Edited identifiers where upper case is used but lower case is required, or vice versa.

Corrected an error in the Number of Sequences field, specifically:

A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.

Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field according due to a PatentIn bug). Sequences corrected:

Other: Number of lines in field 223 exceeded
in 3 of 18 sequences. mtb

*Examiner: The above corrections must be communicated to the applicant in the first C Action. DO NOT send a copy of this form.

RECEIVED

NOV 14 2001

1642

TECH CENTER 1600/2900

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/234,290A

DATE: 11/06/2001
 TIME: 15:54:03

Input Set : N:\jumbos\i234290a.RAW
 Output Set: N:\CRF3\11062001\I234290A.raw

1 <110> APPLICANT: Burkly, Linda C.
 2 <120> TITLE OF INVENTION: TREATMENT FOR INSULIN DEPENDENT DIABETES
 3 <130> FILE REFERENCE: 10274-008003
 C--> 4 <140> CURRENT APPLICATION NUMBER: US/09/234,290A
 5 <141> CURRENT FILING DATE: 1999-01-20
 6 <150> PRIOR APPLICATION NUMBER: US 08/447,118
 7 <151> PRIOR FILING DATE: 1995-05-22
 8 <150> PRIOR APPLICATION NUMBER: US 08/029,330
 9 <151> PRIOR FILING DATE: 1993-02-09
 10 <150> PRIOR APPLICATION NUMBER: PCT/US94/01456
 11 <151> PRIOR FILING DATE: 1994-02-09
 12 <160> NUMBER OF SEQ ID NOS: 18
 13 <170> SOFTWARE: FastSEQ for Windows Version 4.0
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 17 <212> TYPE: DNA
 18 <213> ORGANISM: Homo sapiens
 19 <220> FEATURE:
 20 <221> NAME/KEY: misc_feature
 21 <222> LOCATION: (0)...(0)
 22 <223> OTHER INFORMATION: pBAG159 insert: HP1/2 heavy chain variable region; ✓
 23 amino acid 1 is Glu (E) but Gln (Q) may be
 24 substituted
 25 <221> NAME/KEY: CDS
 26 <222> LOCATION: (1)...(360)
 27 <400> SEQUENCE: 1
 28 gtc aaa ctg cag cag tct ggg gca gag ctt gtg aag cca ggg gcc tca 48
 29 Val Lys Leu Gln Gln Ser Gly Ala Glu Leu Val Lys Pro Gly Ala Ser
 30 1 5 10 15
 31 gtc aag ttg tcc tgc aca gct tct ggc ttc aac att aaa gac acc tat 96
 32 Val Lys Leu Ser Cys Thr Ala Ser Gly Phe Asn Ile Lys Asp Thr Tyr
 33 20 25 30
 34 atg cac tgg gtg aag cag agg cct gaa cag ggc ctg gag tgg att gga 144
 35 Met His Trp Val Lys Gln Arg Pro Glu Gln Gly Leu Glu Trp Ile Gly
 36 35 40 45
 37 agg att gat cct gcg agt ggc gat act aaa tat gac ccg aag ttc cag 192
 38 Arg Ile Asp Pro Ala Ser Gly Asp Thr Lys Tyr Asp Pro Lys Phe Gln
 39 50 55 60
 40 gtc aag gcc act att aca gcg gac acg tcc tcc aac aca gcc tgg ctg 240
 41 Val Lys Ala Thr Ile Thr Ala Asp Thr Ser Ser Asn Thr Ala Trp Leu
 42 65 70 75 80
 43 cag ctc agc agc ctg aca tct gag gac act gcc gtc tac tac tgt gca 288
 44 Gln Leu Ser Ser Leu Thr Ser Glu Asp Thr Ala Val Tyr Tyr Cys Ala
 45 85 90 95
 46 gac gga atg tgg gta tca acg gga tat gct ctg gac ttc tgg ggc caa 336
 47 Asp Gly Met Trp Val Ser Thr Gly Tyr Ala Leu Asp Phe Trp Gly Gln
 48 100 105 110

ENTERED

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/234,290A

DATE: 11/06/2001
TIME: 15:54:03

Input Set : N:\jumbos\i234290a.RAW
Output Set: N:\CRF3\11062001\I234290A.raw

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50 Gly Thr Thr Val Thr Val Ser Ser
51 115 120
53 <210> SEQ ID NO: 2
54 <211> LENGTH: 120
55 <212> TYPE: PRT
56 <213> ORGANISM: Homo sapiens
57 <400> SEQUENCE: 2
58 Val Lys Leu Gln Gln Ser Gly Ala Glu Leu Val Lys Pro Gly Ala Ser 15
59 1 5 10 15
60 Val Lys Leu Ser Cys Thr Ala Ser Gly Phe Asn Ile Lys Asp Thr Tyr 30
61 20 25 30
62 Met His Trp Val Lys Gln Arg Pro Glu Gln Gly Leu Glu Trp Ile Gly 45
63 35 40 45
64 Arg Ile Asp Pro Ala Ser Gly Asp Thr Lys Tyr Asp Pro Lys Phe Gln 60
65 50 55 60
66 Val Lys Ala Thr Ile Thr Ala Asp Thr Ser Ser Asn Thr Ala Trp Leu 80
67 65 70 75
68 Gln Leu Ser Ser Leu Thr Ser Glu Asp Thr Ala Val Tyr Tyr Cys Ala 95
69 85 90 95
70 Asp Gly Met Trp Val Ser Thr Gly Tyr Ala Leu Asp Phe Trp Gly Gln 110
71 100 105 110
72 Gly Thr Thr Val Thr Val Ser Ser
73 115 120
75 <210> SEQ ID NO: 3
76 <211> LENGTH: 318
77 <212> TYPE: DNA
78 <213> ORGANISM: Homo sapiens
79 <220> FEATURE:
80 <221> NAME/KEY: misc_feature
81 <222> LOCATION: (0)...(0)
82 <223> OTHER INFORMATION: pBAG172 insert: HP1/2 light chain variable region
83 <221> NAME/KEY: CDS
84 <222> LOCATION: (1)...(318)
85 <223> OTHER INFORMATION: HP1/2 light chain variable region
86 <400> SEQUENCE: 3
87 agt att gtg atg acc cag act ccc aaa ttc ctg ctt gtt tca gca gga 48
88 Ser Ile Val Met Thr Gln Thr Pro Lys Phe Leu Leu Val Ser Ala Gly
89 1 5 10 15
90 gac agg gtt acc ata acc tgc aag gcc agt cag agt gtg act aat gat 96
91 Asp Arg Val Thr Ile Thr Cys Lys Ala Ser Gln Ser Val Thr Asn Asp
92 20 25 30
93 gta gct tgg tac caa cag aag cca ggg cag tct cct aaa ctg ctg ata 144
94 Val Ala Trp Tyr Gln Gln Lys Pro Gly Gln Ser Pro Lys Leu Leu Ile
95 35 40 45
96 tat tat gca tcc aat cgc tac act gga gtc cct gat cgc ttc act ggc 192
97 Tyr Tyr Ala Ser Asn Arg Tyr Thr Gly Val Pro Asp Arg Phe Thr Gly
98 50 55 60
99 agt gga tat ggg acg gat ttc act ttc acc atc agc act gtg cag gct 240

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/234,290A

DATE: 11/06/2001
TIME: 15:54:03

Input Set : N:\jumbos\i234290a.RAW
Output Set: N:\CRF3\11062001\I234290A.raw

100 Ser Gly Tyr Gly Thr Asp Phe Thr Phe Thr Ile Ser Thr Val Gln Ala
101 65 70 75 80
102 gaa gac ctg gca gtt tat ttc tgt cag cag gat tat agc tct ccg tac 288
103 Glu Asp Leu Ala Val Tyr Phe Cys Gln Gln Asp Tyr Ser Ser Pro Tyr
104 85 90 95
105 acg ttc gga ggg ggg acc aag ctg gag atc 318
106 Thr Phe Gly Gly Thr Lys Leu Glu Ile
107 100 105
109 <210> SEQ ID NO: 4
110 <211> LENGTH: 106
111 <212> TYPE: PRT
112 <213> ORGANISM: Homo sapiens
113 <400> SEQUENCE: 4
114 Ser Ile Val Met Thr Gln Thr Pro Lys Phe Leu Leu Val Ser Ala Gly
115 1 5 10 15
116 Asp Arg Val Thr Ile Thr Cys Lys Ala Ser Gln Ser Val Thr Asn Asp
117 20 25 30
118 Val Ala Trp Tyr Gln Gln Lys Pro Gly Gln Ser Pro Lys Leu Leu Ile
119 35 40 45
120 Tyr Tyr Ala Ser Asn Arg Tyr Thr Gly Val Pro Asp Arg Phe Thr Gly
121 50 55 60
122 Ser Gly Tyr Gly Thr Asp Phe Thr Phe Thr Ile Ser Thr Val Gln Ala
123 65 70 75 80
124 Glu Asp Leu Ala Val Tyr Phe Cys Gln Gln Asp Tyr Ser Ser Pro Tyr
125 85 90 95
126 Thr Phe Gly Gly Thr Lys Leu Glu Ile
127 100 105
129 <210> SEQ ID NO: 5
130 <211> LENGTH: 429
131 <212> TYPE: DNA
132 <213> ORGANISM: Homo sapiens
133 <220> FEATURE:
134 <221> NAME/KEY: CDS
135 <222> LOCATION: (1)...(429)
136 <221> NAME/KEY: sig_peptide
137 <222> LOCATION: (1)...(57)
138 <221> NAME/KEY: mat_peptide
139 <222> LOCATION: (58)...(429)
140 <221> NAME/KEY: misc_feature
141 <222> LOCATION: (0)...(0)
142 <223> OTHER INFORMATION: pBAG195 insert: AS heavy chain variable region
143 <400> SEQUENCE: 5
144 atg gac tgg acc tgg agg gtc ttc tgc ttg ctg gct gta gca cca ggt 48
145 Met Asp Trp Thr Trp Arg Val Phe Cys Leu Leu Ala Val Ala Pro Gly
146 -15 -10 -5
147 gcc cac tcc cag gtc caa ctg cag gag agc ggt cca ggt ctt gtg aga 96
148 Ala His Ser Gln Val Gln Leu Gln Glu Ser Gly Pro Gly Leu Val Arg
149 1 5 10
150 cct agc cag acc ctg agc ctg acc tgc acc gcg tct ggc ttc aac att 144

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/234,290A

DATE: 11/06/2001
TIME: 15:54:03

Input Set : N:\jumbos\i234290a.RAW
Output Set: N:\CRF3\11062001\I234290A.raw

151	Pro Ser Gln Thr Leu Ser Leu Thr Cys Thr Ala Ser Gly Phe Asn Ile			
152	15	20	25	
153	aaa gac acc tat atg cac tgg gtg aga cag cca cct gga cga ggt ctt			192
154	Lys Asp Thr Tyr Met His Trp Val Arg Gln Pro Pro Gly Arg Gly Leu			
155	30	35	40	45
156	gag tgg att gga agg att gat cct gcg agt ggc gat act aaa tat gac			240
157	Glu Trp Ile Gly Arg Ile Asp Pro Ala Ser Gly Asp Thr Lys Tyr Asp			
158	50	55	60	
159	ccg aag ttc cag gtc aga gtg aca atg ctg gta gac acc agc agc aac			288
160	Pro Lys Phe Gln Val Arg Val Thr Met Leu Val Asp Thr Ser Ser Asn			
161	65	70	75	
162	cag ttc agc ctg aga ctc agc agc gtg aca gcc gcc gac acc gcg gtc			336
163	Gln Phe Ser Leu Arg Leu Ser Ser Val Thr Ala Ala Asp Thr Ala Val			
164	80	85	90	
165	tat tat tgt gca gac gga atg tgg gta tca acg gga tat gct ctg gac			384
166	Tyr Tyr Cys Ala Asp Gly Met Trp Val Ser Thr Gly Tyr Ala Leu Asp			
167	95	100	105	
168	ttc tgg ggc caa ggg acc acg gtc acc gtc tcc tca ggt gag tcc			429
169	Phe Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser Gly Glu Ser			
170	110	115	120	
172	<210> SEQ ID NO: 6			
173	<211> LENGTH: 143			
174	<212> TYPE: PRT			
175	<213> ORGANISM: Homo sapiens			
176	<220> FEATURE:			
177	<221> NAME/KEY: SIGNAL			
178	<222> LOCATION: (1)...(19)			
179	<400> SEQUENCE: 6			
180	Met Asp Trp Thr Trp Arg Val Phe Cys Leu Leu Ala Val Ala Pro Gly			
181	-15	-10	-5	
182	Ala His Ser Gln Val Gln Leu Gln Glu Ser Gly Pro Gly Leu Val Arg			
183	1	5	10	
184	Pro Ser Gln Thr Leu Ser Leu Thr Cys Thr Ala Ser Gly Phe Asn Ile			
185	15	20	25	
186	Lys Asp Thr Tyr Met His Trp Val Arg Gln Pro Pro Gly Arg Gly Leu			
187	30	35	40	45
188	Glu Trp Ile Gly Arg Ile Asp Pro Ala Ser Gly Asp Thr Lys Tyr Asp			
189	50	55	60	
190	Pro Lys Phe Gln Val Arg Val Thr Met Leu Val Asp Thr Ser Ser Asn			
191	65	70	75	
192	Gln Phe Ser Leu Arg Leu Ser Ser Val Thr Ala Ala Asp Thr Ala Val			
193	80	85	90	
194	Tyr Tyr Cys Ala Asp Gly Met Trp Val Ser Thr Gly Tyr Ala Leu Asp			
195	95	100	105	
196	Phe Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser Gly Glu Ser			
197	110	115	120	
199	<210> SEQ ID NO: 7			
200	<211> LENGTH: 386			
201	<212> TYPE: DNA			

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/234,290A

DATE: 11/06/2001
TIME: 15:54:03

Input Set : N:\jumbos\i234290a.RAW
Output Set: N:\CRF3\11062001\I234290A.raw

202 <213> ORGANISM: Homo sapiens
 203 <220> FEATURE:
 204 <221> NAME/KEY: CDS
 205 <222> LOCATION: (1)...(384)
 206 <221> NAME/KEY: sig_peptide
 207 <222> LOCATION: (1)...(57)
 208 <221> NAME/KEY: mat_peptide
 209 <222> LOCATION: (58)...(384)
 210 <221> NAME/KEY: misc_feature
 211 <222> LOCATION: (0)...(0)
 212 <223> OTHER INFORMATION: pBAG198 insert: VK2 (SVMDY) light chain variable
 213 region
 214 <400> SEQUENCE: 7
 215 atg ggt tgg tcc tgc atc atc ctg ttc ctg gtt gct acc gct acc ggt 48
 216 Met Gly Trp Ser Cys Ile Ile Leu Phe Leu Val Ala Thr Ala Thr Gly
 217 -15 -10 -5
 218 gtc cac tcc agc atc gtg atg acc cag agc cca agc agc ctg agc gcc 96
 219 Val His Ser Ser Ile Val Met Thr Gln Ser Pro Ser Ser Leu Ser Ala
 220 1 5 10
 221 agc gtg ggt gac aga gtg acc atc acc tgt aag gcc agt cag agt gtg 144
 222 Ser Val Gly Asp Arg Val Thr Ile Thr Cys Lys Ala Ser Gln Ser Val
 223 15 20 25
 224 act aat gat gta gct tgg tac cag cag aag cca ggt aag gct cca aag 192
 225 Thr Asn Asp Val Ala Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys
 226 30 35 40 45
 227 ctg ctg atc tac tat gca tcc aat cgc tac act ggt gtg cca gat aga 240
 228 Leu Leu Ile Tyr Tyr Ala Ser Asn Arg Tyr Thr Gly Val Pro Asp Arg
 229 50 55 60
 230 ttc agc ggt agc ggt tat ggt acc gac ttc acc ttc acc atc agc agc 288
 231 Phe Ser Gly Ser Gly Tyr Gly Thr Asp Phe Thr Phe Thr Ile Ser Ser
 232 65 70 75
 233 ctc cag cca gag gac atc gcc acc tac tac tgc cag cag gat tat agc 336
 234 Leu Gln Pro Glu Asp Ile Ala Thr Tyr Tyr Cys Gln Gln Asp Tyr Ser
 235 80 85 90
 236 tct ccg tac acg ttc ggc caa ggg acc aag gtg gaa atc aaa cgt aag 384
 237 Ser Pro Tyr Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Arg Lys
 238 95 100 105
 239 tg 386
 241 <210> SEQ ID NO: 8
 242 <211> LENGTH: 128
 243 <212> TYPE: PRT
 244 <213> ORGANISM: Homo sapiens
 245 <220> FEATURE:
 246 <221> NAME/KEY: SIGNAL
 247 <222> LOCATION: (1)...(19)
 248 <400> SEQUENCE: 8
 249 Met Gly Trp Ser Cys Ile Ile Leu Phe Leu Val Ala Thr Ala Thr Gly
 250 -15 -10 -5
 251 Val His Ser Ser Ile Val Met Thr Gln Ser Pro Ser Ser Leu Ser Ala

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/234,290A

DATE: 11/06/2001

TIME: 15:54:04

Input Set : N:\jumbos\i234290a.RAW

Output Set: N:\CRF3\11062001\I234290A.raw

L:4 M:270 C: Current Application Number differs, Wrong Format

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/234,290A

DATE: 10/12/2001
TIME: 08:48:11

Input Set : A:\10274-008003.TXT
Output Set: N:\CRF3\10122001\I234290A.raw

4 <110> APPLICANT: Burkly, Linda C.
6 <120> TITLE OF INVENTION: TREATMENT FOR INSULIN DEPENDENT DIABETES
9 <130> FILE REFERENCE: 10274-008003
11 <140> CURRENT APPLICATION NUMBER: US 09/234,290A
12 <141> CURRENT FILING DATE: 1999-01-20
14 <150> PRIOR APPLICATION NUMBER: US 08/447,118
15 <151> PRIOR FILING DATE: 1995-05-22
17 <150> PRIOR APPLICATION NUMBER: US 08/029,330
18 <151> PRIOR FILING DATE: 1993-02-09
20 <150> PRIOR APPLICATION NUMBER: PCT/US94/01456
21 <151> PRIOR FILING DATE: 1994-02-09
23 <160> NUMBER OF SEQ ID NOS: 18
25 <170> SOFTWARE: FastSEQ for Windows Version 4.0
27 <210> SEQ ID NO: 1
28 <211> LENGTH: 360
29 <212> TYPE: DNA
30 <213> ORGANISM: Homo sapiens
32 <220> FEATURE:
33 <221> NAME/KEY: misc_feature
34 <222> LOCATION: (0)...(0)
35 <223> OTHER INFORMATION: pBAG159 insert: HP1/2 heavy chain variable region;
36 amino acid 1 is Glu (E) but Gln (Q) may be
37 substituted
40 <221> NAME/KEY: CDS
41 <222> LOCATION: (1)...(360)
43 <400> SEQUENCE: 1
44 gtc aaa ctg cag cag tct ggg gca gag ctt gtg aag cca ggg gcc tca 48
45 Val Lys Leu Gln Gln Ser Gly Ala Glu Leu Val Lys Pro Gly Ala Ser
46 1 5 10 15
48 gtc aag ttg tcc tgc aca gct tct ggc ttc aac att aaa gac acc tat 96
49 Val Lys Leu Ser Cys Thr Ala Ser Gly Phe Asn Ile Lys Asp Thr Tyr
50 20 25 30
52 atg cac tgg gtg aag cag agg cct gaa cag ggc ctg gag tgg att gga 144
53 Met His Trp Val Lys Gln Arg Pro Glu Gln Gly Leu Glu Trp Ile Gly
54 35 40 45
56 agg att gat cct gcg agt ggc gat act aaa tat gac ccg aag ttc cag 192
57 Arg Ile Asp Pro Ala Ser Gly Asp Thr Lys Tyr Asp Pro Lys Phe Gln
58 50 55 60
60 gtc aag gcc act att aca gcg gac acg tcc tcc aac aca gcc tgg ctg 240
61 Val Lys Ala Thr Ile Thr Ala Asp Thr Ser Ser Asn Thr Ala Trp Leu
62 65 70 75 80
64 cag ctc agc agc ctg aca tct gag gac act gcc gtc tac tac tgt gca 288
65 Gln Leu Ser Ser Leu Thr Ser Glu Asp Thr Ala Val Tyr Tyr Cys Ala
66 85 90 95
68 gac gga atg tgg gta tca acg gga tat gct ctg gac ttc tgg ggc caa 336
69 Asp Gly Met Trp Val Ser Thr Gly Tyr Ala Leu Asp Phe Trp Gly Gln
70 100 105 110

*Edited directly
Field 223 not
4 liner*

Edited raw file

RAW SEQUENCE LISTING DATE: 10/12/2001
 PATENT APPLICATION: US/09/234,290A TIME: 08:48:11

Input Set : A:\10274-008003.TXT
 Output Set: N:\CRF3\10122001\I234290A.raw

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 73 Gly Thr Thr Val Thr Val Ser Ser
 74 115 120
 77 <210> SEQ ID NO: 2
 78 <211> LENGTH: 120
 79 <212> TYPE: PRT
 80 <213> ORGANISM: Homo sapiens
 82 <400> SEQUENCE: 2
 83 Val Lys Leu Gln Gln Ser Gly Ala Glu Leu Val Lys Pro Gly Ala Ser
 84 1 5 10 15
 85 Val Lys Leu Ser Cys Thr Ala Ser Gly Phe Asn Ile Lys Asp Thr Tyr
 86 20 25 30
 87 Met His Trp Val Lys Gln Arg Pro Glu Gln Gly Leu Glu Trp Ile Gly
 88 35 40 45
 89 Arg Ile Asp Pro Ala Ser Gly Asp Thr Lys Tyr Asp Pro Lys Phe Gln
 90 50 55 60
 91 Val Lys Ala Thr Ile Thr Ala Asp Thr Ser Ser Asn Thr Ala Trp Leu
 92 65 70 75 80
 93 Gln Leu Ser Ser Leu Thr Ser Glu Asp Thr Ala Val Tyr Tyr Cys Ala
 94 85 90 95
 95 Asp Gly Met Trp Val Ser Thr Gly Tyr Ala Leu Asp Phe Trp Gly Gln
 96 100 105 110
 97 Gly Thr Thr Val Thr Val Ser Ser
 98 115 120
 100 <210> SEQ ID NO: 3
 101 <211> LENGTH: 318
 102 <212> TYPE: DNA
 103 <213> ORGANISM: Homo sapiens
 105 <220> FEATURE:
 106 <221> NAME/KEY: misc_feature
 107 <222> LOCATION: (0)...(0)
 108 <223> OTHER INFORMATION: pBAG172 insert: HP1/2 light chain variable region
 110 <221> NAME/KEY: CDS
 111 <222> LOCATION: (1)...(318)
 112 <223> OTHER INFORMATION: HP1/2 light chain variable region
 114 <400> SEQUENCE: 3
 115 agt att gtg atg acc cag act ccc aaa ttc ctg ctt gtt tca gca gga 48
 116 Ser Ile Val Met Thr Gln Thr Pro Lys Phe Leu Leu Val Ser Ala Gly
 117 1 5 10 15
 119 gac agg gtt acc ata acc tgc aag gcc agt cag agt gtg act aat gat 96
 120 Asp Arg Val Thr Ile Thr Cys Lys Ala Ser Gln Ser Val Thr Asn Asp
 121 20 25 30
 123 gta gct tgg tac caa cag aag cca ggg cag tct cct aaa ctg ctg ata 144
 124 Val Ala Trp Tyr Gln Gln Lys Pro Gly Gln Ser Pro Lys Leu Leu Ile
 125 35 40 45
 127 tat tat gca tcc aat cgc tac act gga gtc cct gat cgc ttc act ggc 192
 128 Tyr Tyr Ala Ser Asn Arg Tyr Thr Gly Val Pro Asp Arg Phe Thr Gly
 129 50 55 60
 131 agt gga tat ggg acg gat ttc act ttc acc atc agc act gtg cag gct 240

RAW SEQUENCE LISTING DATE: 10/12/2001
 PATENT APPLICATION: US/09/234,290A TIME: 08:48:11

Input Set : A:\10274-008003.TXT
 Output Set: N:\CRF3\10122001\I234290A.raw

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132 Ser Gly Tyr Gly Thr Asp Phe Thr Phe Thr Ile Ser Thr Val Gln Ala
133   65           70           75           80
135 gaa gac ctg gca gtt tat ttc tgt cag cag gat tat agc tct ccg tac      288
136 Glu Asp Leu Ala Val Tyr Phe Cys Gln Gln Asp Tyr Ser Ser Pro Tyr
137           85           90           95
139 acg ttc gga ggg ggg acc aag ctg gag atc      318
140 Thr Phe Gly Gly Thr Lys Leu Glu Ile
141           100          105
144 <210> SEQ ID NO: 4
145 <211> LENGTH: 106
146 <212> TYPE: PRT
147 <213> ORGANISM: Homo sapiens
149 <400> SEQUENCE: 4
150 Ser Ile Val Met Thr Gln Thr Pro Lys Phe Leu Leu Val Ser Ala Gly
151   1           5           10           15
152 Asp Arg Val Thr Ile Thr Cys Lys Ala Ser Gln Ser Val Thr Asn Asp
153           20           25           30
154 Val Ala Trp Tyr Gln Gln Lys Pro Gly Gln Ser Pro Lys Leu Leu Ile
155           35           40           45
156 Tyr Tyr Ala Ser Asn Arg Tyr Thr Gly Val Pro Asp Arg Phe Thr Gly
157           50           55           60
158 Ser Gly Tyr Gly Thr Asp Phe Thr Phe Thr Ile Ser Thr Val Gln Ala
159   65           70           75           80
160 Glu Asp Leu Ala Val Tyr Phe Cys Gln Gln Asp Tyr Ser Ser Pro Tyr
161           85           90           95
162 Thr Phe Gly Gly Thr Lys Leu Glu Ile
163           100          105
165 <210> SEQ ID NO: 5
166 <211> LENGTH: 429
167 <212> TYPE: DNA
168 <213> ORGANISM: Homo sapiens
170 <220> FEATURE:
171 <221> NAME/KEY: CDS
172 <222> LOCATION: (1)...(429)
174 <221> NAME/KEY: sig_peptide
175 <222> LOCATION: (1)...(57)
177 <221> NAME/KEY: mat_peptide
178 <222> LOCATION: (58)...(429)
180 <221> NAME/KEY: misc_feature
181 <222> LOCATION: (0)...(0)
182 <223> OTHER INFORMATION: pBAG195 insert: AS heavy chain variable region
184 <400> SEQUENCE: 5
185 atg gac tgg acc tgg agg gtc ttc tgc ttg ctg gct gta gca cca ggt      48
186 Met Asp Trp Thr Trp Arg Val Phe Cys Leu Leu Ala Val Ala Pro Gly
187           -15          -10          -5
189 gcc cac tcc cag gtc caa ctg cag gag agc ggt cca ggt ctt gtg aga      96
190 Ala His Ser Gln Val Gln Leu Gln Glu Ser Gly Pro Gly Leu Val Arg
191           1           5           10
193 cct agc cag acc ctg agc ctg acc tgc acc gcg tct ggc ttc aac att      144
  
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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/234,290A

DATE: 10/12/2001

TIME: 08:48:11

Input Set : A:\10274-008003.TXT

Output Set: N:\CRF3\10122001\I234290A.raw

194	Pro	Ser	Gln	Thr	Leu	Ser	Leu	Thr	Cys	Thr	Ala	Ser	Gly	Phe	Asn	Ile	
195	15				20				25								
197	aaa	gac	acc	tat	atg	cac	tgg	gtg	aga	cag	cca	cct	gga	cga	ggg	ctt	192
198	Lys	Asp	Thr	Tyr	Met	His	Trp	Val	Arg	Gln	Pro	Pro	Gly	Arg	Gly	Leu	
199	30				35				40				45				
201	gag	tgg	att	gga	agg	att	gat	cct	gcg	agt	ggc	gat	act	aaa	tat	gac	240
202	Glu	Trp	Ile	Gly	Arg	Ile	Asp	Pro	Ala	Ser	Gly	Asp	Thr	Lys	Tyr	Asp	
203						50			55				60				
205	ccg	aag	ttc	cag	gtc	aga	gtg	aca	atg	ctg	gta	gac	acc	agc	agc	aac	288
206	Pro	Lys	Phe	Gln	Val	Arg	Val	Thr	Met	Leu	Val	Asp	Thr	Ser	Ser	Asn	
207						65			70				75				
209	cag	ttc	agc	ctg	aga	ctc	agc	gtg	aca	gcc	gcc	gac	acc	gcg	gtc	336	
210	Gln	Phe	Ser	Leu	Arg	Leu	Ser	Ser	Val	Thr	Ala	Ala	Asp	Thr	Ala	Val	
211						80			85				90				
213	tat	tat	tgt	gca	gac	gga	atg	tgg	gta	tca	acg	gga	tat	gct	ctg	gac	384
214	Tyr	Tyr	Cys	Ala	Asp	Gly	Met	Trp	Val	Ser	Thr	Gly	Tyr	Ala	Leu	Asp	
215						95			100				105				
217	ttc	tgg	ggc	caa	ggg	acc	acg	gtc	acc	gtc	tcc	tca	ggt	gag	tcc	429	
218	Phe	Trp	Gly	Gln	Gly	Thr	Thr	Val	Thr	Val	Ser	Ser	Gly	Glu	Ser		
219						110			115				120				
222	<210> SEQ ID NO: 6																
223	<211> LENGTH: 143																
224	<212> TYPE: PRT																
225	<213> ORGANISM: Homo sapiens																
227	<220> FEATURE:																
228	<221> NAME/KEY: SIGNAL																
229	<222> LOCATION: (1)...(19)																
231	<400> SEQUENCE: 6																
232	Met	Asp	Trp	Thr	Trp	Arg	Val	Phe	Cys	Leu	Leu	Ala	Val	Ala	Pro	Gly	
233					-15				-10				-5				
234	Ala	His	Ser	Gln	Val	Gln	Leu	Gln	Glu	Ser	Gly	Pro	Gly	Leu	Val	Arg	
235					1			5				10					
236	Pro	Ser	Gln	Thr	Leu	Ser	Leu	Thr	Cys	Thr	Ala	Ser	Gly	Phe	Asn	Ile	
237					15			20				25					
238	Lys	Asp	Thr	Tyr	Met	His	Trp	Val	Arg	Gln	Pro	Pro	Gly	Arg	Gly	Leu	
239					30			35			40			45			
240	Glu	Trp	Ile	Gly	Arg	Ile	Asp	Pro	Ala	Ser	Gly	Asp	Thr	Lys	Tyr	Asp	
241					50			55				60					
242	Pro	Lys	Phe	Gln	Val	Arg	Val	Thr	Met	Leu	Val	Asp	Thr	Ser	Ser	Asn	
243					65			70				75					
244	Gln	Phe	Ser	Leu	Arg	Leu	Ser	Ser	Val	Thr	Ala	Ala	Asp	Thr	Ala	Val	
245					80			85				90					
246	Tyr	Tyr	Cys	Ala	Asp	Gly	Met	Trp	Val	Ser	Thr	Gly	Tyr	Ala	Leu	Asp	
247					95			100				105					
248	Phe	Trp	Gly	Gln	Gly	Thr	Thr	Val	Thr	Val	Ser	Ser	Gly	Glu	Ser		
249					110			115				120					
251	<210> SEQ ID NO: 7																
252	<211> LENGTH: 386																
253	<212> TYPE: DNA																

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/234,290A

DATE: 10/12/2001
TIME: 08:48:11

Input Set : A:\10274-008003.TXT
Output Set: N:\CRF3\10122001\I234290A.raw

254 <213> ORGANISM: Homo sapiens
 256 <220> FEATURE:
 257 <221> NAME/KEY: CDS
 258 <222> LOCATION: (1)...(384)
 260 <221> NAME/KEY: sig_peptide
 261 <222> LOCATION: (1)...(57)
 263 <221> NAME/KEY: mat_peptide
 264 <222> LOCATION: (58)...(384)
 266 <221> NAME/KEY: misc_feature
 267 <222> LOCATION: (0)...(0)
 268 <223> OTHER INFORMATION: pBAG198 insert: VK2 (SVMDY) light chain variable
 269 region
 271 <400> SEQUENCE: 7
 272 atg ggt tgg tcc tgc atc atc ctg ttc ctg gtt gct acc gct acc ggt 48
 273 Met Gly Trp Ser Cys Ile Ile Leu Phe Leu Val Ala Thr Ala Thr Gly
 274 -15 -10 -5
 276 gtc cac tcc agc atc gtg atg acc cag agc cca agc agc ctg agc gcc 96
 277 Val His Ser Ser Ile Val Met Thr Gln Ser Pro Ser Ser Leu Ser Ala
 278 1 5 10
 280 agc gtg ggt gac aga gtg acc atc acc tgt aag gcc agt cag agt gtg 144
 281 Ser Val Gly Asp Arg Val Thr Ile Thr Cys Lys Ala Ser Gln Ser Val
 282 15 20 25
 284 act aat gat gta gct tgg tac cag cag aag cca ggt aag gct cca aag 192
 285 Thr Asn Asp Val Ala Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys
 286 30 35 40 45
 288 ctg ctg atc tac tat gca tcc aat cgc tac act ggt gtg cca gat aga 240
 289 Leu Leu Ile Tyr Tyr Ala Ser Asn Arg Tyr Thr Gly Val Pro Asp Arg
 290 50 55 60
 292 ttc agc ggt agc ggt tat ggt acc gac ttc acc ttc acc atc agc agc 288
 293 Phe Ser Gly Ser Gly Tyr Gly Thr Asp Phe Thr Phe Thr Ile Ser Ser
 294 65 70 75
 296 ctc cag cca gag gac atc gcc acc tac tac tgc cag cag gat tat agc 336
 297 Leu Gln Pro Glu Asp Ile Ala Thr Tyr Tyr Cys Gln Gln Asp Tyr Ser
 298 80 85 90
 300 tct ccg tac acg ttc ggc caa ggg acc aag gtg gaa atc aaa cgt aag 384
 301 Ser Pro Tyr Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Arg Lys
 302 95 100 105
 304 tg 386
 306 <210> SEQ ID NO: 8
 307 <211> LENGTH: 128
 308 <212> TYPE: PRT
 309 <213> ORGANISM: Homo sapiens
 311 <220> FEATURE:
 312 <221> NAME/KEY: SIGNAL
 313 <222> LOCATION: (1)...(19)
 315 <400> SEQUENCE: 8
 316 Met Gly Trp Ser Cys Ile Ile Leu Phe Leu Val Ala Thr Ala Thr Gly
 317 -15 -10 -5
 318 Val His Ser Ser Ile Val Met Thr Gln Ser Pro Ser Ser Leu Ser Ala

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/234,290A

DATE: 10/12/2001

TIME: 08:48:12

Input Set : A:\10274-008003.TXT

Output Set: N:\CRF3\10122001\I234290A.raw

L:355 M:259 W: Allowed number of lines exceeded, <223> Other Information:

L:363 M:259 W: Allowed number of lines exceeded, <223> Other Information:

L:371 M:259 W: Allowed number of lines exceeded, <223> Other Information:

STATISTICS SUMMARY

PATENT APPLICATION: US/09/234,290A

DATE: 10/12/2001

TIME: 08:48:12

Input Set : A:\10274-008003.TXT

Output Set: N:\CRF3\10122001\I234290A.raw

Application Serial Number: US/09/234,290A

Alpha or Numeric: Numeric

Application Class:

Application File Date: 01-20-1999

Art Unit: 1642

Software Application: FastSeq

Total Number of Sequences: 18

Total Nucleotides: 3081

Total Amino Acids: 956

Number of Errors: 0

Number of Warnings: 3

Number of Corrections: 0

MESSAGE SUMMARY

259 W: 3 (Allowed number of lines exceeded)